

# PCT 371 Routing Sheet

## APPLICATION

10/526,028

IFW DocCode- SEQREQ

Index using Current Date

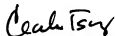
### TO BE DELIVERED TO:

Tech Center Scanning

### Sequence Rule Compliance Review Item

<b>X</b>	CRF, paper copy of sequence listing, and statement that both are same missing
	CRF contains error(s) according to STIC Report
	CRF damaged or unreadable according to STIC Report
	CRF transferred from prior application is not compliant
	OTHER

**Place an "X" in the appropriate box**



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Supervisory Patent Examiner/  
Technology Center 1000

# APPLICATION SERIAL NUMBER

10/526,028

## DOES NOT COMPLY WITH THE SEQUENCE RULES. See reasons below.

**Claim 7 recites a sequence embraced by the sequence rules, however, Applicant has not provided a CRF, paper copy or statement that they are the same, as required.**

37 CFR § 1.821 (a) states,

“Nucleotide and/or amino acid sequences as used in §§ 1.821 through 1.825 are interpreted to mean an unbranched sequence of four or more amino acids or an unbranched sequence of ten or more nucleotides. Branched sequences are specifically excluded from this definition. Sequences with fewer than four specifically defined nucleotides or amino acids are specifically excluded from this section. “Specifically defined” means those amino acids other than “Xaa” and those nucleotide bases other than “n” defined in accordance with the World Intellectual Property Organization (WIPO) Handbook on Industrial Property Information and Documentation, Standard ST.25: Standard for the Presentation of Nucleotide and Amino Acid Sequence Listings in Patent Applications (1998), including Tables 1 through 6 in Appendix 2.”

(a)(2) states,

“*Amino acids*: Amino acids are those L-amino acids commonly found in naturally occurring proteins and are listed in WIPO Standard ST.25 (1998), Appendix 2, Table 3. Those amino acid sequences containing D-amino acids are not intended to be embraced by this definition. Any amino acid sequence that contains post-translationally modified amino acids may be described as the amino acid sequence that is initially translated using the symbols shown in WIPO Standard ST.25 (1998), Appendix 2, Table 3 with the modified positions; e.g., hydroxylations or glycosylations, being described as set forth in WIPO Standard ST.25 (1998), Appendix 2, Table 4, but these modifications shall not be shown explicitly in the amino acid sequence. Any peptide or protein that can be expressed as a sequence using the symbols in WIPO Standard ST.25 (1998), Appendix 2, Table 3 in conjunction with a description in the Feature section to describe, for example, modified linkages, cross links and end caps, non-peptidyl bonds, etc., is embraced by this definition.”